

SEQUENCE LISTING

<110> Blaschuk, Orest W.
Symonds, James Matthew
Gour, Barbara J.

<120> COMPOUNDS AND METHODS FOR MODULATING NONCLASSICAL
CADHERIN-MEDIATED FUNCTIONS

<130> 100086.407C7

<140> US
<141> 2001-12-03

<160> 4052

<170> PatentIn Ver. 2.0

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Motif in Extracellular domains of Classical
Cadherins

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<223> Where Xaa is Isoleucine, Leucine or Valine

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<223> Where Xaa is Aspartic Acid, Asparagine or Glutamic Acid

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<213> Homo sapiens

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Arg Ser Lys Arg Gly Trp Val Trp Asn Gln Phe Phe Val Ile Glu Glu
1 5 10 15

Tyr Thr Gly Pro Asp Pro Val Leu Val Gly Arg Leu His Ser Asp Ile
20 25 30

Asp Ser Gly Asp Gly Asn Ile Lys Tyr Ile Leu Ser Gly Glu Gly Ala
35 40 45

Gly Thr Ile Phe Val Ile Asp Asp Lys Ser Gly Asn Ile His Ala Thr
50 55 60

Lys Thr Leu Asp Arg Glu Glu Arg Ala Gln Tyr Thr Leu Met Ala Gln

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Lys	Asn	Thr	Ser	Leu	Pro	His	His	Val	Gly	Lys	Ile	Lys	Ser	Ser	Val	
			20					25					30			
Ser	Arg	Lys	Asn	Ala	Lys	Tyr	Leu	Leu	Lys	Gly	Glu	Tyr	Val	Gly	Lys	
		35				40						45				
Val	Phe	Arg	Val	Asp	Ala	Glu	Thr	Gly	Asp	Val	Phe	Ala	Ile	Glu	Arg	
	50					55					60					
Leu	Asp	Arg	Glu	Asn	Ile	Ser	Glu	Tyr	His	Leu	Thr	Ala	Val	Ile	Val	
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Asp	Lys	Asp	Thr	Gly	Glu	Asn	Leu	Glu	Thr	Pro	Ser	Ser	Phe	Thr	Ile	
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 Asp Arg Gly Asp Gly Ser Leu Lys Tyr Ile Leu Ser Gly Asp Gly Ala
 35 40 45
 Gly Asp Leu Phe Ile Ile Asn Glu Asn Thr Gly Asp Ile Gln Ala Thr
 50 55 60
 Lys Arg Leu Asp Arg Glu Glu Lys Pro Val Tyr Ile Leu Arg Ala Gln
 65 70 75 80
 Ala Ile Asn Arg Arg Thr Gly Arg Pro Val Glu Pro Glu Ser Glu Phe
 85 90 95
 Ile Ile Lys Ile His Asp Ile Asn Asp Asn Glu Pro Ile Phe
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 Thr Lys Glu Val Tyr Thr Ala Thr Val Pro Glu Met Ser Asp Val Gly
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 Gly Asn Ser Ala Lys Val Val Tyr Ser Ile Leu Gln Gly Gln Pro Tyr
 35 40 45
 Phe Ser Val Glu Ser Glu Thr Gly Ile Ile Lys Thr Ala Leu Leu Asn
 50 55 60
 Met Asp Arg Glu Asn Arg Glu Gln Tyr Gln Val Val Ile Gln Ala Lys
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 Asp Met Gly Gly Gln Met Gly Gly Leu Ser Gly Thr Thr Thr Val Asn
 85 90 95
 Ile Thr Leu Thr Asp Val Asn Asp Asn Pro Pro Arg Phe
 100 105

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Leu Asp Gly Pro Tyr Thr Ala Gly Val Pro Glu Met Ser Pro Val Gly
 1 5 10 15
 Thr Ser Val Val Gln Val Thr Ala Thr Asp Ala Asp Asp Pro Thr Tyr
 20 25 30
 Gly Asn Ser Ala Arg Val Val Tyr Ser Ile Leu Gln Gly Gln Pro Tyr
 35 40 45
 Phe Ser Val Glu Pro Lys Thr Gly Ile Ile Lys Thr Ala Leu Pro Asn
 50 55 60
 Met Asp Arg Glu Ala Lys Asp Gln Tyr Leu Leu Val Ile Gln Ala Lys
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 85 90 95
 Val Thr Leu Thr Asp Val Asn Asp Asn Pro Pro Arg Phe
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<210> 12
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 <213> Gallus gallus

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 20 25 30
 Pro Val Arg Tyr Ser Ile Asp Arg Asn Thr Asp Leu Glu Arg Tyr Phe
 35 40 45
 Asn Ile Asp Ala Asn Ser Gly Val Ile Thr Thr Ala Lys Ser Leu Asp
 50 55 60
 Arg Glu Thr Asn Ala Val His Asn Ile Thr Val Leu Ala Met Glu Ser
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 85 90 95
 Asp Ile Asn Asp Asn Ala Pro Glu Phe
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<400> 13
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 Phe Ser Gly Pro Glu Pro Ile Leu Val Gly Arg Leu His Thr Asp Leu
 20 25 30

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Asp Pro Gly Ser Lys Lys Ile Lys Tyr Ile Leu Ser Gly Asp Gly Ala
 35 40 45
 Gly Thr Ile Phe Gln Ile Asn Asp Val Thr Gly Asp Ile His Ala Ile
 50 55 60
 Lys Arg Leu Asp Arg Glu Lys Ala Glu Tyr Thr Leu Thr Ala Gln
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 Ala Val Asp Trp Glu Thr Ser Lys Pro Leu Glu Pro Pro Ser Glu Phe
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 Ile Ile Lys Val Gln Asp Ile Asn Asp Asn Ala Pro Glu Phe
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 Arg Val Lys Arg Gly Trp Val Trp Asn Gln Phe Phe Val Leu Glu Glu
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 Tyr Val Gly Ser Glu Pro Gln Tyr Val Gly Lys Leu His Ser Asp Leu
 20 25 30
 Asp Lys Gly Glu Gly Thr Val Lys Tyr Thr Leu Ser Gly Asp Gly Ala
 35 40 45
 Gly Thr Val Phe Thr Ile Asp Glu Thr Thr Gly Asp Ile His Ala Ile
 50 55 60
 Arg Ser Leu Asp Arg Glu Glu Lys Pro Phe Tyr Thr Leu Arg Ala Gln
 65 70 75 80
 Ala Val Asp Ile Glu Thr Arg Lys Pro Leu Glu Pro Glu Ser Glu Phe
 85 90 95
 Ile Ile Lys Val Gln Asp Ile Asn Asp Asn Glu Pro Lys Phe
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<210> 15
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<400> 15
 Leu Asp Gly Pro Tyr Val Ala Thr Val Pro Glu Met Ser Pro Val Gly
 1 5 10 15
 Ala Tyr Val Leu Gln Val Lys Ala Thr Asp Ala Asp Asp Pro Thr Tyr
 20 25 30
 Gly Asn Ser Ala Arg Val Val Tyr Ser Ile Leu Gln Gly Gln Pro Tyr
 35 40 45
 Phe Ser Ile Asp Pro Lys Thr Gly Val Ile Arg Thr Ala Leu Pro Asn

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<210> 17
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<400> 17
Thr Asp Gly Pro Tyr Ile Val Thr Val Pro Glu Met Ser Asp Met Gly
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Thr Ser Val Leu Gln Val Thr Ala Thr Asp Ala Asp Asp Pro Thr Tyr
      20      25      30
Gly Asn Ser Ala Arg Val Val Tyr Ser Ile Leu Gln Gly Gln Pro Tyr
      35      40      45
Phe Ser Val Asp Pro Lys Thr Gly Val Ile Arg Thr Ala Leu His Asn
      50      55      60
Met Asp Arg Glu Ala Arg Glu His Tyr Ser Val Val Ile Gln Ala Lys
      65      70      75      80

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Ile Thr Leu Thr Asp Val Asn Asp Asn Pro Pro Arg Phe
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Ser	Met	Pro	Ser	Tyr	Leu	Met	Glu	Val	Tyr	Glu	Asn	Ala	Lys	Ile	Gly	
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Thr	Val	Val	Gly	Thr	Val	Leu	Ala	Gln	Asp	Pro	Asp	Ser	Thr	Asn	Ser	
			20					25					30			
Leu	Val	Arg	Tyr	Phe	Ile	Asn	Tyr	Asn	Val	Glu	Asp	Asp	Arg	Phe	Phe	
		35					40					45				
Asn	Ile	Asp	Ala	Asn	Thr	Gly	Thr	Ile	Arg	Thr	Thr	Lys	Val	Leu	Asp	
50						55					60					
Arg	Glu	Glu	Thr	Pro	Trp	Tyr	Asn	Ile	Thr	Val	Thr	Ala	Ser	Glu	Ile	
65					70					75					80	
Asp	Asn	Pro	Asp	Leu	Leu	Ser	His	Val	Thr	Val	Gly	Ile	Arg	Val	Leu	
				85					90					95		
Asp	Val	Asn	Asp	Asn	Pro	Pro	Glu	Leu								
			100					105								

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<210> 19
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<400> 19															
Arg	Val	Arg	Arg	Ala	Trp	Val	Ile	Pro	Pro	Ile	Ser	Val	Ser	Glu	Asn
1				5					10					15	
His	Lys	Arg	Leu	Pro	Tyr	Pro	Leu	Val	Gln	Ile	Lys	Ser	Asp	Lys	Gln
			20					25					30		
Gln	Leu	Gly	Ser	Val	Ile	Tyr	Ser	Ile	Gln	Gly	Pro	Gly	Val	Asp	Glu
		35					40					45			
Glu	Pro	Arg	Gly	Val	Phe	Ser	Ile	Asp	Lys	Phe	Thr	Gly	Lys	Val	Phe
	50					55					60				
Leu	Asn	Ala	Met	Leu	Asp	Arg	Glu	Lys	Thr	Asp	Arg	Phe	Arg	Leu	Arg
65					70					75					80
Ala	Phe	Ala	Leu	Asp	Leu	Gly	Gly	Ser	Thr	Leu	Glu	Asp	Pro	Thr	Asp
				85					90					95	
Leu	Glu	Ile	Val	Val	Val	Asp	Gln	Asn	Asp	Asn	Arg	Pro	Ala	Phe	
			100					105					110		

<210> 20
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<400> 20
 Leu Gln Glu Ala Phe Thr Gly Arg Val Leu Glu Gly Ala Val Pro Gly
 1 5 10 15
 Thr Tyr Val Thr Arg Ala Glu Ala Thr Asp Ala Asp Asp Pro Glu Thr
 20 25 30
 Asp Asn Ala Ala Leu Arg Phe Ser Ile Leu Gln Gln Gly Ser Pro Glu
 35 40 45
 Leu Phe Ser Ile Asp Glu Leu Thr Gly Glu Ile Arg Thr Val Gln Val
 50 55 60
 Gly Leu Asp Arg Glu Val Val Ala Val Tyr Asn Leu Thr Leu Gln Val
 65 70 75 80
 Ala Asp Met Ser Gly Asp Gly Leu Thr Ala Thr Ala Ser Ala Ile Ile
 85 90 95
 Thr Leu Asp Asp Ile Asn Asp Asn Ala Pro Glu Phe
 100 105

<210> 21
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<400> 21
 Arg Gln Lys Arg Ser Ile Val Val Ser Pro Ile Leu Ile Pro Glu Asn
 1 5 10 15
 Gln Arg Gln Pro Phe Pro Arg Asp Val Gly Lys Val Val Asp Ser Asp
 20 25 30
 Arg Pro Glu Arg Ser Lys Phe Arg Leu Thr Gly Lys Gly Val Asp Gln
 35 40 45
 Glu Pro Lys Gly Ile Phe Arg Ile Asn Glu Asn Thr Gly Ser Val Ser
 50 55 60
 Val Thr Arg Thr Leu Asp Arg Glu Val Ile Ala Val Tyr Gln Leu Phe
 65 70 75 80
 Val Glu Thr Thr Asp Val Asn Gly Lys Thr Leu Glu Gly Pro Val Pro
 85 90 95
 Leu Glu Val Ile Val Ile Asp Gln Asn Asp Asn Arg Pro Ile Phe
 100 105 110

<210> 22
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<213> Rattus norvegicus

<400> 22

Arg Val Lys Arg Gly Trp Val Trp Asn Gln Phe Phe Val Val Glu Glu
1 5 10 15

Tyr Thr Gly Thr Glu Pro Leu Tyr Val Gly Lys Ile His Ser Asp Ser
20 25 30

Asp Glu Gly Asp Gly Thr Ile Lys Tyr Thr Ile Ser Gly Glu Gly Ala
35 40 45

Gly Thr Ile Phe Leu Ile Asp Glu Leu Thr Gly Asp Ile His Ala Thr
50 55 60

Glu Arg Leu Asp Arg Glu Gln Lys Thr Phe Tyr Thr Leu Arg Ala Gln
65 70 75 80

Ala Arg Asp Arg Ala Thr Asn Arg Leu Leu Glu Pro Glu Ser Glu Phe
85 90 95

Ile Ile Lys Val Gln Asp Ile Asn Asp Ser Glu Pro Arg Phe
100 105 110

<210> 23

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<213> Rattus norvegicus

<400> 23

Leu His Gly Pro Tyr Ile Gly Ser Val Ala Glu Leu Ser Pro Thr Gly
1 5 10 15

Thr Ser Val Met Gln Val Met Ala Ser Asp Ala Asp Asp Pro Thr Tyr
20 25 30

Gly Ser Ser Ala Arg Leu Val Tyr Ser Val Leu Asp Gly Glu His His
35 40 45

Phe Thr Val Asp Pro Lys Thr Gly Val Ile Arg Thr Ala Val Pro Asp
50 55 60

Leu Asp Arg Glu Ser Gln Glu Arg Tyr Glu Val Val Ile Gln Ala Thr
65 70 75 80

Asp Met Ala Gly Gln Leu Gly Gly Leu Ser Gly Ser Thr Thr Val Thr
85 90 95

Ile Val Val Thr Asp Val Asn Asp Asn Pro Pro Arg Phe
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<213> Rattus norvegicus

<400> 24

Arg Pro Pro Ser Gly Leu Leu Glu Val Gln Glu Asp Ala Gln Val Gly
1 5 10 15

Ser Leu Val Gly Val Val Thr Ala Arg Asp Pro Asp Ala Ala Asn Arg
 20 25 30
 Pro Val Arg Tyr Ala Ile Asp Arg Asp Ser Asp Leu Glu Gln Ile Phe
 35 40 45
 Asp Ile Asp Ala Asp Thr Gly Ala Ile Val Thr Gly Lys Gly Leu Asp
 50 55 60
 Arg Glu Thr Ala Gly Trp His Asn Ile Thr Val Leu Ala Met Glu Ala
 65 70 75 80
 Asp Asn His Ala Gln Leu Ser Arg Ala Ser Leu Arg Ile Arg Ile Leu
 85 90 95
 Asp Val Asn Asp Asn Pro Pro Glu Leu
 100 105

<210> 25
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<400> 25
 Leu Gln Ser Lys Tyr Glu Gly Ser Val Arg Gln Asn Ser Arg Pro Gly
 1 5 10 15
 Lys Pro Phe Leu Tyr Val Asn Ala Thr Asp Leu Asp Asp Pro Ala Thr
 20 25 30
 Pro Asn Gly Gln Leu Tyr Tyr Gln Ile Val Ile Gln Leu Pro Met Ile
 35 40 45
 Asn Asn Val Met Tyr Phe Gln Ile Asn Asn Lys Thr Gly Ala Ile Ser
 50 55 60
 Leu Thr Arg Glu Gly Ser Gln Glu Leu Asn Pro Ala Lys Asn Pro Tyr
 65 70 75 80
 Asn Leu Val Ile Ser Val Lys Asp Met Gly Gly Gln Ser Glu Asn Ser
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 Lys Ala Pro Ala Pro
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 Arg Leu Asp Phe Glu Asp Thr Lys Leu His Glu Ile Tyr Ile Gln Ala
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Glu Leu Phe Ser Ile Asp Pro Lys Thr Gly Leu Ile Arg Val Lys Gly
 50 55 60
 Asn Leu Asp Tyr Glu Glu Asn Gly Met Leu Glu Ile Asp Val Gln Ala
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 35 40 45
 Asp Asp His Leu Phe Glu Ile Asp Pro Ser Ser Gly Glu Ile Arg Thr
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 35 40 45
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 35 40 45
 Ser Pro Met Phe Ile Ile Asn Arg Asn Thr Gly Glu Ile Arg Thr Met
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 65 70 75 80
 Arg Gly Ser Asp Arg Asp Gly Gly Ala Asp Gly Met Ser Ala Glu Cys
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<210> 32
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 35 40 45
 Pro Pro Val Phe Tyr Leu Asn Lys Asp Thr Gly Glu Ile Tyr Thr Thr
 50 55 60
 Ser Val Thr Leu Asp Arg Glu Glu His Ser Ser Tyr Thr Leu Thr Val
 65 70 75 80
 Glu Ala Arg Asp Gly Asn Gly Glu Val Thr Asp Lys Pro Val Lys Gln

85 90 95
 Ala Gln Val Gln Ile Arg Ile Leu Asp Val Asn Asp Asn Ile Pro Val
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Val

<210> 33
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<400> 33
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 Phe Pro Gln His Val Gln Gln Ile Gln Ser Asp Ala Ala Gln Asn Tyr
 20 25 30
 Thr Ile Phe Tyr Ser Ile Ser Gly Pro Gly Val Asp Lys Glu Pro Phe
 35 40 45
 Asn Leu Phe Tyr Ile Glu Lys Asp Thr Gly Asp Ile Phe Cys Thr Arg
 50 55 60
 Ser Ile Asp Arg Glu Lys Tyr Glu Gln Phe Ala Leu Tyr Gly Tyr Ala
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 85 90 95
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<210> 34
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<400> 34
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 Thr Ile Tyr Tyr Ser Ile Arg Gly Pro Gly Val Asp Gln Glu Pro Arg
 35 40 45
 Asn Leu Phe Tyr Val Glu Arg Asp Thr Gly Asn Leu Tyr Cys Thr Arg
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 Pro Val Asp Arg Glu Gln Tyr Glu Ser Phe Glu Ile Ile Ala Phe Ala
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Lys Ile Glu Asp Glu Asn Asp Asn Tyr Pro Ile
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<210> 35
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<400> 35
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Thr Val Phe Tyr Ser Ile Ser Gly Arg Gly Val Asp Lys Glu Pro Leu
35 40 45
Asn Leu Phe Tyr Ile Glu Arg Asp Thr Gly Asn Leu Phe Cys Thr Arg
50 55 60
Pro Val Asp Arg Glu Glu Tyr Asp Val Phe Asp Leu Ile Ala Tyr Ala
65 70 75 80
Ser Thr Ala Asp Gly Tyr Ser Ala Asp Leu Pro Leu Pro Leu Pro Ile
85 90 95
Arg Val Glu Asp Glu Asn Asp Asn His Pro Val
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<210> 36
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<400> 36
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Thr Val Phe Tyr Ser Ile Ser Gly Arg Gly Val Asp Lys Glu Pro Leu
35 40 45
Asn Leu Phe Tyr Ile Glu Arg Asp Thr Gly Asn Leu Phe Cys Thr Arg
50 55 60
Pro Val Asp Arg Glu Glu Tyr Asp Val Phe Asp Leu Ile Ala Tyr Ala
65 70 75 80
Ser Thr Ala Asp Gly Tyr Ser Ala Asp Leu Pro Leu Pro Leu Pro Ile
85 90 95
Arg Val Glu Asp Glu Asn Asp Asn His Pro Val
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<210> 37

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 <213> Mus musculus

<400> 37
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 Thr Leu Val Val Lys Val Asn Ala Ser Asp Leu Asp Glu Gly Val Asn
 20 25 30
 Gly Asp Ile Met Tyr Ser Phe Ser Thr Asp Ile Ser Pro Asn Val Lys
 35 40 45
 Tyr Lys Phe His Ile Asp Pro Val Ser Gly Glu Ile Ile Val Lys Gly
 50 55 60
 Tyr Ile Asp Phe Glu Glu Cys Lys Ser Tyr Glu Ile Leu Ile Glu Gly
 65 70 75 80
 Ile Asp Lys Gly Gln Leu Pro Leu Ser Gly His Cys Lys Val Ile Val
 85 90 95
 Gln Val Glu Asp Ile Asn Asp Asn Val Pro Glu Leu
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<210> 38
 <211> 108
 <212> PRT
 <213> Mus musculus

<400> 38
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 Thr Thr Val Ile Arg Leu Asn Ala Ser Asp Lys Asp Glu Gly Thr Asn
 20 25 30
 Ser Ala Ile Ser Tyr Ser Phe Asn Arg Leu Val Pro Pro Lys Thr Leu
 35 40 45
 Glu Gln Phe Ser Ile Asp Ala Asp Thr Gly Glu Ile Ile Thr Gln Gly
 50 55 60
 Asn Leu Asp Phe Glu Gln Val Asp Val Tyr Lys Ile His Val Asp Ala
 65 70 75 80
 Thr Asp Lys Gly His Pro Pro Met Val Gly His Cys Thr Val Leu Val
 85 90 95
 Lys Val Leu Asp Glu Asn Asp Asn Val Pro Gln Ile
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<210> 39
 <211> 108
 <212> PRT
 <213> Mus musculus

<400> 39

Asp Arg Ala Ile Tyr Arg Val Lys Leu Val Glu Asn Ala Arg Asn Gly
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 Thr Val Val Ile Arg Leu Asn Ala Ser Asp Leu Asp Glu Gly Ser Asn
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 Gly Gln Ile Leu Tyr Ser Phe Ala Ala Asp Val Ser Pro Lys Thr Glu
 35 40 45
 Ala Thr Phe His Ile Asp Ser Val Ser Gly Glu Ile Lys Val Asn Gly
 50 55 60
 Lys Ile Asp Phe Glu Glu Thr Asn Leu Trp Lys Ile Gln Ala Glu Ala
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 Val Asp Lys Gly Ser Pro Pro Met Phe Gly His Cys Thr Ile Leu Ile
 85 90 95
 Glu Val Leu Asp Ile Asn Asp Asn Ala Pro Lys Ile
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<210> 40
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 <213> Mus musculus

<400> 40
 Asp Arg Phe Val Tyr Lys Val Lys Val Leu Glu Asp Ala Leu Asn Gly
 1 5 10 15
 Thr Leu Val Ile Asn Leu Asn Ala Thr Asp Pro Asp Glu Gly Ile Asn
 20 25 30
 Gly Asp Ile Ile Tyr Ser Phe Arg Arg Pro Val Ser Pro Ala Val Val
 35 40 45
 His Ala Phe Asn Ile Asp Ser Asn Ser Gly Glu Val Arg Thr Lys Gly
 50 55 60
 Leu Leu Asp Phe Glu Glu Ile Lys Leu Tyr Glu Ile Pro Val Glu Ala
 65 70 75 80
 Val Asp Lys Gly Asn Ile Pro Met Thr Gly His Cys Thr Leu Leu Val
 85 90 95
 Glu Leu Leu Asp Val Asn Asp Asn Ala Pro Glu Val
 100 105

<210> 41
 <211> 108
 <212> PRT
 <213> Mus musculus

<400> 41
 Asp Lys Ser Ile Tyr Asn Val Arg Leu Leu Glu Asn Thr Pro Asn Gly
 1 5 10 15
 Thr Leu Val Ile Lys Leu Asn Ala Ser Asp Ala Asp Glu Gly Ile Asn
 20 25 30

Lys Glu Ile Leu Tyr Phe Phe Ser Asn Leu Val Leu Asp Asp Val Lys
 35 40 45
 Ser Lys Phe Thr Ile Asp Ser Ser Ser Gly Glu Ile Lys Val Lys Gly
 50 55 60
 Glu Leu Asp Tyr Glu Asp Cys Lys Val Tyr Glu Ile Asn Ile Asp Ala
 65 70 75 80
 Val Asp Arg Ser Ala Phe Pro Leu Ala Gly His Cys Lys Ile Ile Val
 85 90 95
 Lys Leu Val Asp Val Asn Asp Asn Val Pro Glu Met
 100 105

<210> 42
 <211> 107
 <212> PRT
 <213> Mus musculus

<400> 42
 Asp His Leu Glu Tyr Lys Val Arg Ile Met Glu Asn Ala Ala Lys Glu
 1 5 10 15
 Thr Leu Val Ile Thr Leu Asn Ala Thr Asp Leu Asp Glu Gly Ala Asn
 20 25 30
 Gly Gln Leu Val Tyr Ser Leu Met Ser Ile Lys Pro Thr Gly Arg His
 35 40 45
 Leu Phe Thr Leu Asp Glu Lys Asn Gly Glu Leu Arg Val Asn Gly Thr
 50 55 60
 Leu Asp Tyr Glu Glu Asn Lys Leu Tyr Glu Ile Glu Val Leu Ala Thr
 65 70 75 80
 Asp Lys Gly Thr Pro Pro Met Val Gly His Cys Val Val Leu Val Glu
 85 90 95
 Ile Leu Asp Thr Asn Asp Asn Ser Pro Glu Val
 100 105

<210> 43
 <211> 108
 <212> PRT
 <213> Mus musculus

<400> 43
 Asp Arg Ser Val Tyr Glu Val Lys Met Tyr Glu Asn Gln Glu Asn Lys
 1 5 10 15
 Thr Leu Val Ile Trp Leu Asn Ala Thr Asp Ser Asp Glu Gly Ile Asn
 20 25 30
 Lys Glu Val Glu Tyr Ser Phe Ser Ser Leu Ala Ser Ser Ile Ile Arg
 35 40 45
 Gln Lys Phe Leu Ile Asn Glu Lys Thr Gly Glu Ile Lys Ile Asn Gly

50 55 60
 Ala Ile Asp Phe Glu Glu Ser Asn Asn Tyr Glu Ile His Val Asp Ala
 65 70 75 80
 Thr Asp Lys Gly Tyr Pro Pro Met Val Ala His Cys Thr Val Leu Val
 85 90 95
 Glu Ile Leu Asp Glu Asn Asp Asn Ala Pro Glu Ile
 100 105

<210> 44
 <211> 106
 <212> PRT
 <213> Homo sapiens

<400> 44
 Gly Trp Val Trp Asn Gln Phe Phe Val Ile Glu Glu Tyr Thr Gly Pro
 1 5 10 15
 Asp Pro Val Leu Val Gly Arg Leu His Ser Asp Ile Asp Ser Gly Asp
 20 25 30
 Gly Asn Ile Lys Tyr Ile Leu Ser Gly Glu Gly Ala Gly Thr Ile Phe
 35 40 45
 Val Ile Asp Asp Lys Ser Gly Asn Ile His Ala Thr Lys Thr Leu Asp
 50 55 60
 Arg Glu Glu Arg Ala Gln Tyr Thr Leu Met Ala Gln Ala Val Asp Arg
 65 70 75 80
 Asp Thr Asn Arg Pro Leu Glu Pro Pro Ser Glu Phe Ile Val Lys Val
 85 90 95
 Gln Asp Ile Asn Asp Asn Pro Pro Glu Phe
 100 105

<210> 45
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<400> 45
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 Asp Pro Val Leu Val Gly Arg Leu His Ser Asp Ile Asp Ser Gly Asp
 20 25 30
 Gly Asn Ile Lys Tyr Ile Leu Ser Gly Glu Gly Ala Gly Thr Ile Phe
 35 40 45
 Val Ile Asp Asp Lys Ser Gly Asn Ile His Ala Thr Lys Thr Leu Asp
 50 55 60
 Arg Glu Glu Arg Ala Gln Tyr Thr Leu Met Ala Gln Ala Val Asp Arg
 65 70 75 80

Asp Thr Asn Arg Pro Leu Glu Pro Pro Ser Glu Phe Ile Val Lys Val
 85 90 95

Gln Asp Ile Asn Asp Asn Pro Pro Glu Phe
 100 105

<210> 46
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 Motifs in the Extracellular Domains of
 Nonclassical Cadherins

<220>
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 <222> (1)
 <223> Where Xaa is any amino acid

<220>
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 <222> (3)
 <223> Where Xaa is any amino acid

<400> 46
 Xaa Asp Xaa Glu
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<210> 47
 <211> 5
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Calcium Binding
 Motifs in the Extracellular Domains of
 Nonclassical Cadherins

<220>
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 <223> Where Xaa is any amino acid

<220>
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 <222> (3)
 <223> Where Xaa is any amino acid

<220>
 <221> VARIANT
 <222> (5)
 <223> Where Xaa is any amino acid

<400> 47
 Asp Xaa Xaa Asp Xaa
 1 5

<210> 48
 <211> 5
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Classical
 Cadherin Cell Adhesion Recognition Sequence

<400> 48
 Tyr Ile Gly Ser Arg
 1 5

<210> 49
 <211> 10
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Classical
 Cadherin Cell Adhesion Recognition Sequence

<400> 49
 Lys Tyr Ser Phe Asn Tyr Asp Gly Ser Glu
 1 5 10

<210> 50
 <211> 17
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: N-CAM Heparin
 Sulfate-Binding Site

<400> 50
 Ile Trp Lys His Lys Gly Arg Asp Val Ile Leu Lys Lys Asp Val Arg
 1 5 10 15

Phe

<210> 51
 <211> 4
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Putative Claudin
 Cell Adhesion Recognition Sequence

<400> 51
 Ile Tyr Ser Tyr
 1

<210> 52

<211> 4
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Occludin Cell
 Adhesion Recognition Sequence

<400> 52
 Leu Tyr His Tyr
 1

<210> 53
 <211> 10
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<220>
 <223> Cyclic Peptide

<220>
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 <222> (2)
 <223> Where Xaa is beta,beta-tetramethylene cysteine

<220>
 <223> Description of Artificial Sequence: Product of
 Synthesis and Cyclization based on Human
 OB-Cadherin

<400> 53
 Ile Xaa Val Ile Asp Asp Lys Ser Cys Glu
 1 5 10

<210> 54
 <211> 10
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Product of
 Synthesis and Cyclization based on Human
 OB-Cadherin

<220>
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<220>
 <221> VARIANT
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<400> 54
 Ile Xaa Val Ile Asp Asp Lys Ser Gly Cys
 1 5 10

<210> 55
 <211> 9

<212> PRT
 <213> Artificial Sequence

<220>
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 Synthesis and Cyclization based on Human
 OB-Cadherin

<220>
 <223> Cyclic Peptide

<220>
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 <222> (1)
 <223> Where Xaa is beta-mercaptopropionic acid

<400> 55
 Xaa Val Ile Asp Asp Lys Ser Gly Cys
 1 5

<210> 56
 <211> 9
 <212> PRT
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<220>
 <223> Description of Artificial Sequence: Product of
 Synthesis and Cyclization based on Human
 OB-Cadherin

<220>
 <223> Cyclic Peptide

<220>
 <221> VARIANT
 <222> (1)
 <223> Where Xaa is
 beta,beta-pentamethylene-beta-mercaptopropionic
 acid

<400> 56
 Xaa Val Ile Asp Asp Lys Ser Gly Cys
 1 5

<210> 57
 <211> 5
 <212> PRT
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<220>
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<220>
 <221> VARIANT
 <222> (4)
 <223> Where Xaa is D-Serine

<220>
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Synthesis and Cyclization based on Human
OB-Cadherin

<400> 57

Asp Asp Lys Xaa Ser
1 5

<210> 58

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Product of
Synthesis and Cyclization based on Human
OB-Cadherin

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<223> Cyclic Peptide

<400> 58

Trp Gly Gly Trp
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<210> 59

<211> 6

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: E-Cadherin Cell
Adhesion Recognition Sequence

<400> 59

Ser His Ala Val Ser Ser
1 5

<210> 60

<211> 6

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: N-Cadherin Cell
Adhesion Recognition Sequence

<400> 60

Ala His Ala Val Asp Ile
1 5

<210> 61

<211> 15

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: N-Cadherin Cell

Adhesion Recognition Sequence

<400> 61

Phe His Leu Arg Ala His Ala Val Asp Ile Asn Gly Asn Gln Val
 1 5 10 15

<210> 62

<211> 48

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: Occludin Cell
 Adhesion Recognition Sequence

<400> 62

Gly Val Asn Pro Thr Ala Gln Ser Ser Gly Ser Leu Tyr Gly Ser Gln
 1 5 10 15

Ile Tyr Ala Leu Cys Asn Gln Phe Tyr Thr Pro Ala Ala Thr Gly Leu
 20 25 30

Tyr Val Asp Gln Tyr Leu Tyr His Tyr Cys Val Val Asp Pro Gln Glu
 35 40 45

<210> 63

<211> 10

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: N-CAM Cell
 Adhesion Recognition Sequence

<400> 63

Lys Tyr Ser Phe Asn Tyr Asp Gly Ser Glu
 1 5 10

<210> 64

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Product of
 Synthesis based on Human Cadherin-5 Cell Adhesion
 Recognition Sequence

<400> 64

Val Phe Arg Val Asp Ala Glu Thr Gly Asp
 1 5 10

<210> 65

<211> 4

<212> PRT
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<220>
<223> Description of Unknown Organism: Calcium Binding
Motifs in the Extracellular Domains of
Nonclassical Cadherins

<400> 65
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<210> 66
<211> 4
<212> PRT
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<220>
<223> Description of Unknown Organism: Calcium Binding
Motifs in the Extracellular Domains of
Nonclassical Cadherins

<400> 66
Leu Asp Phe Glu
1

<210> 67
<211> 4
<212> PRT
<213> Unknown

<220>
<223> Description of Unknown Organism: Calcium Binding
Motifs in the Extracellular Domains of
Nonclassical Cadherins

<400> 67
Leu Asp Tyr Glu
1

<210> 68
<211> 4
<212> PRT
<213> Unknown

<220>
<223> Description of Unknown Organism: Calcium Binding
Motifs in the Extracellular Domains of
Nonclassical Cadherins

<400> 68
Ile Asp Arg Glu
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<210> 69
<211> 4
<212> PRT

$\langle 220 \rangle$

<400> 69

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$\langle 211 \rangle$ 4

<213> Unknown

 $\langle 220 \rangle$

<400> 70

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<211> 5

<213> Artificial Sequence

 $\langle 220 \rangle$

<221> VARIANT

<222> (5)

<223> Where Xaa is beta,beta-dimethyl cysteine

 $\langle 220 \rangle$

<223> Cyclic Peptide

 $\langle 220 \rangle$

<223> Description of Artificial Sequence: Product of
Synthesis and Cyclization based on Human
OB-cadherin

<400> 71

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5

<210> 72

<211> 11

<212> PRT

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<220> Description of Unknown Organism: Calcium Binding
<221> Motifs in Extracellular Domains of Nonclassical
<222> Cadherins

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 <222> (10)
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 Xaa Xaa Xaa Xaa Asp Xaa Asn Asp Xaa Xaa Pro
 1 5 10

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<220>
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<400> 73
 Asp Asp Lys Ser
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<210> 74
 <211> 5
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 OB-cadherin cell adhesion recognition sequence

<400> 74
Val Ile Asp Asp Lys
1 5

<210> 75
<211> 5
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OB-cadherin cell adhesion recognition sequence

<400> 75
Ile Asp Asp Lys Ser
1 5

<210> 76
<211> 6
<212> PRT
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OB-cadherin cell adhesion recognition sequence

<400> 76
Val Ile Asp Asp Lys Ser
1 5

<210> 77
<211> 5
<212> PRT
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<220>
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OB-cadherin cell adhesion recognition sequence

<400> 77
Asp Asp Lys Ser Gly
1 5

<210> 78
<211> 6
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<220>
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OB-cadherin cell adhesion recognition sequence

<400> 78
Ile Asp Asp Lys Ser Gly
1 5

<210> 79

<211> 7
 <212> PRT
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<220>
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 OB-cadherin cell adhesion recognition sequence

<400> 79
 Val Ile Asp Asp Lys Ser Gly
 1 5

<210> 80
 <211> 6
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<220>
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 OB-cadherin cell adhesion recognition sequence

<400> 80
 Phe Val Ile Asp Asp Lys
 1 5

<210> 81
 <211> 7
 <212> PRT
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 OB-cadherin cell adhesion recognition sequence

<400> 81
 Phe Val Ile Asp Asp Lys Ser
 1 5

<210> 82
 <211> 8
 <212> PRT
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<400> 82
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 1 5

<210> 83
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OB-cadherin cell adhesion recognition sequence

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1 5

<210> 84

<211> 8

<212> PRT

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<400> 84

Ile Phe Val Ile Asp Asp Lys Ser
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<210> 85

<211> 9

<212> PRT

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OB-cadherin cell adhesion recognition sequence

<400> 85

Ile Phe Val Ile Asp Asp Lys Ser Gly
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<210> 86

<211> 4

<212> PRT

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OB-cadherin cell adhesion recognition sequence

<400> 86

Ile Glu Glu Tyr
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<210> 87

<211> 4

<212> PRT

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OB-cadherin cell adhesion recognition sequence

<400> 87

Glu Glu Tyr Thr

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<210> 88
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<400> 88

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<212> PRT
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<400> 89

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<211> 6
<212> PRT
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<400> 90

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<210> 91
<211> 5
<212> PRT
<213> Artificial Sequence
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<400> 91

<210>	92
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OB-cadherin cell adhesion recognition sequence

<400> 92

Ile Glu Glu Tyr Thr Gly
1 5

<210> 93

<211> 7

<212> PRT

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OB-cadherin cell adhesion recognition sequence

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Val Ile Glu Glu Tyr Thr Gly
1 5

<210> 94

<211> 6

<212> PRT

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OB-cadherin cell adhesion recognition sequence

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Phe Val Ile Glu Glu Tyr
1 5

<210> 95

<211> 7

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OB-cadherin cell adhesion recognition sequence

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<210> 96

<211> 8

<212> PRT

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OB-cadherin cell adhesion recognition sequence

<400> 96
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<210> 97
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 <212> PRT
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 OB-cadherin cell adhesion recognition sequence

<400> 97
 Phe Phe Val Ile Glu Glu Tyr
 1 5

<210> 98
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
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 OB-cadherin cell adhesion recognition sequence

<400> 98
 Phe Phe Val Ile Glu Glu Tyr Thr
 1 5

<210> 99
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
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 OB-cadherin cell adhesion recognition sequence

<400> 99
 Phe Phe Val Ile Glu Glu Tyr Thr Gly
 1 5

<210> 100
 <211> 4
 <212> PRT
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<220>
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<400> 100
 Val Glu Ala Gln
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<210> 101
<211> 4
<212> PRT
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<220>
<223> Representative linear modulating agent based on
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<400> 101
Glu Ala Gln Thr
1

<210> 102
<211> 5
<212> PRT
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<220>
<223> Representative linear modulating agent based on
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<400> 102
Ser Val Glu Ala Gln
1             5

<210> 103
<211> 5
<212> PRT
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<220>
<223> Representative linear modulating agent based on
      OB-cadherin cell adhesion recognition sequence

<400> 103
Val Glu Ala Gln Thr
1             5

<210> 104
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
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      OB-cadherin cell adhesion recognition sequence

<400> 104
Ser Val Glu Ala Gln Thr
1             5

<210> 105
<211> 5
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<220>

<223> Representative linear modulating agent based on
OB-cadherin cell adhesion recognition sequence

<400> 105

Glu Ala Gln Thr Gly
1 5

<210> 106

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

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OB-cadherin cell adhesion recognition sequence

<400> 106

Val Glu Ala Gln Thr Gly
1 5

<210> 107

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Representative linear modulating agent based on
OB-cadherin cell adhesion recognition sequence

<400> 107

Ser Val Glu Ala Gln Thr Gly
1 5

<210> 108

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Representative linear modulating agent based on
OB-cadherin cell adhesion recognition sequence

<400> 108

Phe Ser Val Glu Ala Gln
1 5

<210> 109

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Representative linear modulating agent based on
OB-cadherin cell adhesion recognition sequence

<400> 109

Phe Ser Val Glu Ala Gln Thr
1 5

<210> 110
<211> 8
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OB-cadherin cell adhesion recognition sequence

<400> 110

Phe Ser Val Glu Ala Gln Thr Gly
1 5

<210> 111
<211> 7
<212> PRT
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<220>
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OB-cadherin cell adhesion recognition sequence

<400> 111

Tyr Phe Ser Val Glu Ala Gln
1 5

<210> 112
<211> 8
<212> PRT
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<220>
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OB-cadherin cell adhesion recognition sequence

<400> 112

Tyr Phe Ser Val Glu Ala Gln Thr
1 5

<210> 113
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
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OB-cadherin cell adhesion recognition sequence

<400> 113

Tyr Phe Ser Val Glu Ala Gln Thr Gly
1 5

<210> 114

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<211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Representative linear modulating agent based on
 cadherin-5 cell adhesion recognition sequence

<400> 114
 Val Asp Ala Glu
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<210> 115
 <211> 4
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<220>
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 cadherin-5 cell adhesion recognition sequence

<400> 115
 Asp Ala Glu Thr
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<210> 116
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
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 cadherin-5 cell adhesion recognition sequence

<400> 116
 Arg Val Asp Ala Glu
 1 5

<210> 117
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
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 cadherin-5 cell adhesion recognition sequence

<400> 117
 Val Asp Ala Glu Thr
 1 5

<210> 118
 <211> 6
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Representative linear modulating agent based on
cadherin-5 cell adhesion recognition sequence

<400> 118
Arg Val Asp Ala Glu Thr
1 5

<210> 119
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Representative linear modulating agent based on
cadherin-5 cell adhesion recognition sequence

<400> 119
Asp Ala Glu Thr Gly
1 5

<210> 120
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
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cadherin-5 cell adhesion recognition sequence

<400> 120
Val Asp Ala Glu Thr Gly
1 5

<210> 121
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Representative linear modulating agent based on
cadherin-5 cell adhesion recognition sequence

<400> 121
Arg Val Asp Ala Glu Thr Gly
1 5

<210> 122
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
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cadherin-5 cell adhesion recognition sequence

<400> 122
Phe Arg Val Asp Ala Glu

1

5

<210> 123
 <211> 7
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<220>
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 cadherin-5 cell adhesion recognition sequence

<400> 123

Phe Arg Val Asp Ala Glu Thr

1

5

<210> 124
 <211> 8
 <212> PRT
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<220>
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 cadherin-5 cell adhesion recognition sequence

<400> 124

Phe Arg Val Asp Ala Glu Thr Gly

1

5

<210> 125
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<220>
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 cadherin-5 cell adhesion recognition sequence

<400> 125

Val Phe Arg Val Asp Ala Glu

1

5

<210> 126
 <211> 8
 <212> PRT
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<220>
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 cadherin-5 cell adhesion recognition sequence

<400> 126

Val Phe Arg Val Asp Ala Glu Thr

1

5

<210> 127
 <211> 9
 <212> PRT

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<220>

<223> Representative linear modulating agent based on
cadherin-5 cell adhesion recognition sequence

<400> 127

Val Phe Arg Val Asp Ala Glu Thr Gly
1 5

<210> 128

<211> 4

<212> PRT

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<220>

<223> Representative linear modulating agent based on
cadherin-6 cell adhesion recognition sequence

<400> 128

Ile Asn Glu Asn
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<210> 129

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Representative linear modulating agent based on
cadherin-6 cell adhesion recognition sequence

<400> 129

Asn Glu Asn Thr
1

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OB-cadherin cell adhesion recognition sequence

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OB-cadherin cell adhesion recognition sequence

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OB-cadherin cell adhesion recognition sequence

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Protocadherin cell adhesion recognition sequence

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PRT = Proline-rich type

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Patent application of the University of California, San Diego

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Desmocollin

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Cys Ile Asp Ser Asn Ser Gly Cys
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Lys Gln Phe Ser Ile Asp Ala Asp Asp
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cadherin-related neuronal receptor cell adhesion
recognition sequence

Lys Gln Phe Ser Ile Asp Ala Asp Thr Asp
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cadherin-related neuronal receptor cell adhesion
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Lys Gln Phe Ser Ile Asp Ala Asp Thr Gly Asp
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Lys Asp Ser Val Asp
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Lys His Ile Asp Ser Val Asp
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Lys His Ile Asp Ser Val Ser Asp
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Lys His Ile Asp Ser Val Ser Gly Asp
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Lys Asp Ser Ser Asp
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Lys Asp Glu Lys Asn Gly Asp
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Lys Ile Asn Glu Lys Asp
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Glu Ile Asn Glu Asn Thr Gly Lys
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